CRF E████s Corrected by th STIC System█ranch

Serial Number: ___C9/937,2C4_____ 5000

CRF Processing Date: ___5|1|02____
Edited by: ___DC___
Verified by: ___`\DC`___ (STIC staff)

☐ Changed a file from non-ASCII to ASCII

☐ Changed the margins in cases where th sequence text was "wrapped" down to the next line.

☐ Edited a format error in the Current Application Data section, specifically: ENTERED
_____

☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.

☐ Added the mandatory heading and subheadings for "Current Application Data".

☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically:
_____

☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
_____

☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
_____

☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐ Inserted colons after headings/subheadings. Headings edited included:
_____

☐ Deleted extra, invalid, headings used by an applicant, specifically:
_____

☑ Deleted: ☑ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as_____ .

☐ Inserted mandatory headings, specifically: _____

☐ Corrected an obvious error in the response, specifically:
_____

☐ Edited identifiers where upper case is used but lower case is required, or vice versa.

☐ Corrected an error in the Number of Sequences field, specifically:
_____

☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

☐ Other:
_____
_____
_____

*Examiner: The abov corrections must be communicated to the applicant in th first Offic Action. DO NOT send a copy of this form.
                                                                            3/1/95

PCT09

RAW SEQUENCE LISTING   DATE: 05/01/2002
PATENT APPLICATION: US/09/937,204   TIME: 11:41:47

Input Set : A:\PTO.DC.txt
Output Set: N:\CRF3\05012002\I937204.raw

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 4 <110> APPLICANT: Fritig, Bernard
 5        Toquin, Valerie
 6        Geoffroy, Pierrette
 7        Legrand, Michel
 8        Kauffmann, Serge
11 <120> TITLE OF INVENTION: INDUCIBLE COMTII PROMOTER, CHIMERA GENE
12        CONTAINING SAME AND TRANSFORMED PLANTS
15 <130> FILE REFERENCE: A34638-PCT-USA
17 <140> CURRENT APPLICATION NUMBER: US 09/937,204
18 <141> CURRENT FILING DATE: 2000-03-22
20 <150> PRIOR APPLICATION NUMBER: PCT/FR99/03700
21 <151> PRIOR FILING DATE: 1999-03-22
23 <150> PRIOR APPLICATION NUMBER: PCT/FR99/07646
24 <151> PRIOR FILING DATE: 1999-06-11
26 <160> NUMBER OF SEQ ID NOS: 26
28 <170> SOFTWARE: FastSEQ for Windows Version 4.0
30 <210> SEQ ID NO: 1
31 <211> LENGTH: 1863
32 <212> TYPE: DNA
33 <213> ORGANISM: Nicotiana tabacum
35 <220> FEATURE:
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37 <222> LOCATION: (667)...(672)
38 <223> OTHER INFORMATION: Inverted W box
40 <221> NAME/KEY: misc_RNA
41 <222> LOCATION: (820)...(830)
42 <223> OTHER INFORMATION: Inverted L box
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```
45 <222> LOCATION: (845)...(852)
```
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```
48 <222> LOCATION: (1034)...(1047)
49 <223> OTHER INFORMATION: P box
```
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```
52 <222> LOCATION: (1221)...(1226)
53 <223> OTHER INFORMATION: G box
```
W--> 55 <221> misc_signal
```
56 <222> LOCATION: (1343)...(1356)
57 <223> OTHER INFORMATION: Inverse L box
```
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60 <222> LOCATION: (1369)...(1374)
61 <223> OTHER INFORMATION: A box
```
W--> 63 <221> misc_signal
```
64 <222> LOCATION: (1377)...(1382)
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RAW SEQUENCE LISTING   DATE: 05/01/2002
PATENT APPLICATION: US/09/937,204   TIME: 11:41:47

Input Set : A:\PTO.DC.txt
Output Set: N:\CRF3\05012002\I937204.raw

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        68 <222> LOCATION: (1483)...(1488)
        69 <223> OTHER INFORMATION: GT box
.W--> 71 <221> misc_signal
        72 <222> LOCATION: (1562)...(1567)
        73 <223> OTHER INFORMATION: Inverted W box
.W--> 75 <221> misc_signal
        76 <222> LOCATION: (1600)...(1614)
        77 <223> OTHER INFORMATION: L box
W--> 79 <221> CAAT_signal
        80 <222> LOCATION: (1675)...(1679)
W--> 82 <221> misc_signal
        83 <222> LOCATION: (1681)...(1690)
        84 <223> OTHER INFORMATION: E box
W--> 86 <221> CAAT_signal
        87 <222> LOCATION: (1695)...(1699)
W--> 89 <221> TATA_signal
   J    90 <222> LOCATION: (1735)...(1739)
W--> 92 <221> misc_signal
        93 <222> LOCATION: (1772)...(0)
        94 <223> OTHER INFORMATION: Transcription origin
W--> 96 <400> 1
        97 aaagttaggg acaatctata gtgtcacaaa gttgcttatg gcttttggtt cagataaaga 60
        98 aaaagaacag cattttaatt tgtgaagatt agtctgagca gaatttcatt gtatctagaa 120
        99 agaaattgaa aaaagaaata ttctatttca ctattatgtt aggtgcaact atatcatcac 180
       100 catggaaaag ccggagtaaa aagagaacgt agaggagatt tcatgatttg attgagaata 240
       101 taatatatta ttttttttgta attccacaca aagattaaga aaatgatctg atcaatgatg 300
       102 gctccgagga tttggctgtc gcgggaacta tgacattaat ataaatttgt cgctgcctat 360
       103 aaagacccta tctatctatc tatctatcta tatatatata tatatatata tatatatata 420
       104 tatatatata tatatatata tatatatata tatatataag cgctaatatt tgattatttt 480
       105 ttaaaaatat ttataagtat atatgaaatt tttgacgaaa tttttgtgtg accgtgaccc 540
       106 ctcaacctat agtgtgcgtc cacctgtgcc aacaatatag agacaatttg ctcgtatagt 600
       107 cagaaagagt gttttacttt ttagttgctt tttagtgaat ctactcggta taaagttaaa 660
       108 ttagtgggtc aataagtcgg gtgaatagtt aaagaaaaca gtggtgagtt tagctgtcaa 720
       109 ataatttctt cttttttcttg ttttcacatt agaaatcaaa ataaaacaca agcttttttgt 780
       110 atttattta acacaagcta attatatgtt tatatgctgg ttaggtgaag taaagcatgt 840
       111 tatatgagga aagtacgaag aaaatgtgcc aattgtcgtg tacagcaaag cagccagcac 900
       112 aagcaaattc gcacttgata agtggctaag tccactttct agtggaccta gtggttcact 960
       113 aactttttacc aaaaaggcaa taatttgcaa ttcaaaaaga aaaaggaaa aagaaaact 1020
       114 agacagactt taacacacca actcccacag gaagcaacaa tgcaactcac aaaaggaaac 1080
       115 cgagttttttc cgcgacggat ctagaatttg ggttcattct ttacgctttt tcgtattaaa 1140
       116 ctcattatat ttgtataatt atgggtttat attttttatt tattgtaatt tttgtaaaat 1200
       117 tttatatata agtgtatact ccacgtctcc ggatactaca ttagcctcta gggttcttaa 1260
       118 tactcttgtt aaattgtcca ggctccaaac gcatgttcgt ttcaatttta acggatgttt 1320
       119 ccgaacaact ccaaatgttc aatgttaggt gtgtttggtg ttaagcttcc gtcctaggtt 1380
       120 aatagaatag ataattgttg tttcttatat agttttgaac aatcgtcgcc ataaactaat 1440
       121 ttttaggatg gaagctaatt tttaggatgg agtacagcct aaggttaaaa tataactata 1500
       122 aaaaatatcc ataaaaggtg aaatttaatt agtaacatga aaagataaaa ctagtgttat 1560
```

RAW SEQUENCE LISTING   DATE: 05/01/2002
PATENT APPLICATION: US/09/937,204   TIME: 11:41:47

Input Set : A:\PTO.DC.txt
Output Set: N:\CRF3\05012002\I937204.raw

```
123 cggtcaaact ttcaaaagag aaagaaataa ctagacaaac ttcaacaacc aacctgccca 1620
124 acatgctact gtgcaattga aaaataaaca aaagagaacc agacaatatt tcaaccaata 1680
125 ttccatcaag aaaaccaatt atgacaattc ttaaccaaag tcacaactaa cacttataaa 1740
126 aagcactaac tcaactgtac atgattgtga agcctaacaa aaacactcta aaaggaaaag 1800
127 actacgagaa taattacact acaactctta tagctaattc ttgtctcaag attttcagct 1860
128 atg                                                               1863
130 <210> SEQ ID NO: 2
131 <211> LENGTH: 5371
132 <212> TYPE: DNA
133 <213> ORGANISM: Nicotiana tabacum
135 <220> FEATURE:
136 <221> NAME/KEY: promoter
137 <222> LOCATION: (1)...(1860)
139 <221> NAME/KEY: misc_signal
140 <222> LOCATION: (1772)...(0)
141 <223> OTHER INFORMATION: Transcription origin
```
W--> 143 <221> intron
```
144 <222> LOCATION: (2282)...(3633)
```
W--> 146 <221> intron
```
147 <222> LOCATION: (3945)...(4726)
```
W--> 149 <221> terminator
```
150 <222> LOCATION: (5090)...(5371)
```
W--> 152 <400> 2
```
153 aaagttaggg acaatctata gtgtcacaaa gttgcttatg gcttttggtt cagataaaga 60
154 aaaagaacag cattttaatt tgtgaagatt agtctgagca gaatttcatt gtatctagaa 120
155 agaaattgaa aaaagaaata ttctatttca ctattatgtt aggtgcaact atatcatcac 180
156 catggaaaag ccggagtaaa aagagaacgt agaggagatt tcatgatttg attgagaata 240
157 taatatatta ttttttttgta attccacaca aagattaaga aaatgatctg atcaatgatg 300
158 gctccgagga tttggctgtc gcgggaacta tgacattaat ataaatttgt cgctgcctat 360
159 aaagaccta tctatctatc tatctatcta tatatatata tatatatata tatatatata 420
160 tatatatata tatatatata tatatatata tatatataag cgctaatatt tgattatttt 480
161 ttaaaaatat ttataagtat atatgaaatt tttgacgaaa tttttgtgtg accgtgaccc 540
162 ctcaacctat agtgtgcgtc cacctgtgcc aacaatatag agacaatttg ctcgtatagt 600
163 cagaaagagt gttttacttt ttagttgctt tttagtgaat ctactcggta taaagttaaa 660
164 ttagtgggtc aataagtcgg gtgaatagtt aaagaaaaca gtggtgagtt tagctgtcaa 720
165 ataatttctt ctttttcttg ttttcacatt agaaatcaaa ataaaacaca agctttttgt 780
166 atttatttta acacaagcta attatatgtt tatatgctgg ttaggtgaag taaagcatgt 840
167 tatatgagga aagtacgaag aaaatgtgcc aattgtcgtg tacagcaaag cagccagcac 900
168 aagcaaattc gcacttgata agtggctaag tccactttct agtggaccta gtggttcact 960
169 aacttttacc aaaaaggcaa taatttgcaa ttcaaaaga aaaaaggaaa aagaaaact 1020
170 agacagactt taacacacca actcccacag gaagcaacaa tgcaactcac aaaaggaaac 1080
171 cgagtttttc cgcgacggat ctagaatttg ggttcattct ttacgctttt tcgtattaaa 1140
172 ctcattatat ttgtataatt atgggtttat atttttttatt tattgtaatt tttgtaaaat 1200
173 tttatatata agtgtatact ccacgtctcc ggatactaca ttagcctcta gggttcttaa 1260
174 tactcttgtt aaattgtcca ggctccaaac gcatgttcgt ttcaatttta acggatgttt 1320
175 ccgaacaact ccaaatgttc aatgttaggt gtgtttggtg ttaagcttcc gtcctaggtt 1380
176 aatagaatag ataattgttg tttcttatat agttttgaac aatcgtcgcc ataaactaat 1440
177 ttttaggatg gaagctaatt tttaggatgg agtacagcct aaggttaaaa tataactata 1500
178 aaaaatatcc ataaaaggtg aaatttaatt agtaacatga aaagataaaa ctagtgttat 1560
```

RAW SEQUENCE LISTING   DATE: 05/01/2002
PATENT APPLICATION: US/09/937,204   TIME: 11:41:47

Input Set : A:\PTO.DC.txt
Output Set: N:\CRF3\05012002\I937204.raw

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179 cggtcaaact ttcaaaagag aaagaaataa ctagacaaac ttcaacaacc aacctgccca 1620
180 acatgctact gtgcaattga aaaataaaca aaagagaacc agacaatatt tcaaccaata 1680
181 ttccatcaag aaaaccaatt atgacaattc ttaaccaaag tcacaactaa cacttataaa 1740
182 aagcactaac tcaactgtac atgattgtga agcctaacaa aaacactcta aaaggaaaag 1800
183 actacgagaa taattacact acaactctta tagctaattc ttgtctcaag attttcagct 1860
184 atggaatcct caaccaaaag ccaaatacca acacaatcag aagaagagcg taactgcaca 1920
185 tatgccatgc aactattgtc atcttcagtc ctcccctttg tgttgcattc aacaattcaa 1980
186 ttggaagttt ttgagatatt agccaaatct aatgacacta aactttctgc ttctcaaatt 2040
187 gtttctcaaa ttcctaactg cacaaaacct gaagcaccta ctatgttaaa taggatgctt 2100
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189 gggggccaaa aaagagtgta tggtttgtca caagtgggaa aattctttgt taaaaatgaa 2220
190 aatggtgcat caatggggcc acttttggct ttgcttcaaa ataaagtatt cataaacagc 2280
191 tggtaagttt tgtcctactg tgtattcttt ttgcagtggc tgtattgatt ggttgccttt 2340
192 ttcacaagac aagattctta agttttatta cttgtcgatt tatgttagtc gtatgtgcta 2400
193 gtgttattat tctccatctg atcctttat tggtcacttt acctaaaaat attgttacaa 2460
194 aacatttgtc cttctagaaa atcaggtatt attaattttt caattccatc tttattactc 2520
195 caatagtgaa tatggttatt aattagtgtt ttaaggaaga tgtaaggata atttaatcaa 2580
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197 atcttaatca gagtttgatg ttcgagccac aggaatgaat ttgttttaa tagggagtat 2700
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199 tgcgcggggt tcggggaaag gaccgcatca ccaggtctat tgtacgcagc gttacccaac 2820
200 gtgaatctaa attaatgaga ctaaaaaatg gaacccaaca ccagtgaaaa ccaaaaaaag 2880
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205 tacaactaat gttttgagaa aacaaaataa caacaacatc aaacccaatg aaatcccaca 3180
206 agtagagttt ggggaggata gtgtgtacgg agaccttacc cctaccttat aaagttaaag 3240
207 aggctgtttt cgaaagactc tcggctcaag aacattaaaa atttgagaaa acaaaatata 3300
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211 aagtgtcaca ttgttggaga tgggggcaac gctcacaaca aagacgattc tataattagt 3540
212 gttcgaacct gaaattttag ttaaagataa agaagtactt accataatgg tagatatgat 3600
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216 aaaaaatact tgaaaattac aaaggttttg agaaccttaa aactttggtt gatgttggag 3840
217 gtggtcttgg agttaacctc aagatgatta catctaaata ccccacaatt aagggcacta 3900
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219 cttattttgt tgtttattat atttacttcg atcatcaggt ctaggtctgt caagttaaat 4020
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221 agttgtatgc tctgagtctt gcgcatggta tcatagtcaa tttatttaag ctcgttattg 4140
222 cacttgtgaa ttctattata taaggagtaa gcctaccaaa aaggagcgaa aatattttcc 4200
223 aaaactcttt ttaaaccttc ctcaccccat tcccctctcc cctctccccc aacaccaccc 4260
224 accaccccaa ctcccccgtc ttagtttttt tatttatcct ggactttctt atatttatg 4320
225 ctttcctta attgaactct tgtaactaaa ccattgccc cccaccctat agtgtttgcc 4380
226 taaatttat attttttcaaa ataatatttt ctatttacta attaaacatt agaaaatatt 4440
227 tttcggattt ttttccactc accaaccaag catgggaaaa tagtgataaa actactcatt 4500
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228 tttcaaaata atattttcaa ggaaaacatt ttcctttata ccaaataccc ttactcttgt 4560
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230 aaccacatga acaaaatggt tgagctctgc gaattgtgat atatgatttg cttatgtgtt 4680
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233 tcatgactgg agtgatagtc acaacctcaa gttgctaaag aactgctaca aggctctacc 4860
234 agacaatgga aaggtgattg ttgttgaggc cattttacca gtgaaaccag acattgacac 4920
235 cgcagtggtt ggcgtttcgc aatgtgattt gatcatgatg gctcaaaatc ctggaggcaa 4980
236 agagcgatcg gaagaggagt ttcgagcctt ggctactgaa gctggattca aaggcgttaa 5040
237 cttaatatgt tgtgtctgta atttttgggt catggaattc tgcaagtaga tttctactgt 5100
238 acattgagtt tctactactc ttgagtatcc atttatggca atctgggact ggaattgcag 5160
239 cttagtccag attgaacatt gatattccta ataatatttc tattatttcc cttgtttatt 5220
240 tctcttgtat gaaaggatgt cattttgagt attgataatc atgttctcta ggacagaaat 5280
241 tgtaactttg tccaacttta ttgatattcc tagtaagatt tatatgacat gtgtctctgg 5340
242 tttgagaaga gtttcaatat ctacagacgg g                                  5371
243 <210> SEQ ID NO: 3
244 <211> LENGTH: 1095
245 <212> TYPE: DNA
246 <213> ORGANISM: Nicotiana tabacum
248 <220> FEATURE:
249 <221> NAME/KEY: CDS
250 <222> LOCATION: (1)...(1095)
252 <400> SEQUENCE: 3
253 atg gaa tcc tca acc aaa agc caa ata cca aca caa tca gaa gaa gag   48
254 Met Glu Ser Ser Thr Lys Ser Gln Ile Pro Thr Gln Ser Glu Glu Glu
255 1               5                   10                  15
257 cgt aac tgc aca tat gcc atg caa cta ttg tca tct tca gtc ctc ccc   96
258 Arg Asn Cys Thr Tyr Ala Met Gln Leu Leu Ser Ser Ser Val Leu Pro
259             20                  25                  30
261 ttt gtg ttg cat tca aca att caa ttg gaa gtt ttt gag ata tta gcc  144
262 Phe Val Leu His Ser Thr Ile Gln Leu Glu Val Phe Glu Ile Leu Ala
263         35                  40                  45
265 aaa tct aat gac act aaa ctt tct gct tct caa att gtt tct caa att  192
266 Lys Ser Asn Asp Thr Lys Leu Ser Ala Ser Gln Ile Val Ser Gln Ile
267     50                  55                  60
269 cct aac tgc aca aaa cct gaa gca cct act atg tta aat agg atg ctt  240
270 Pro Asn Cys Thr Lys Pro Glu Ala Pro Thr Met Leu Asn Arg Met Leu
271 65                  70                  75                  80
273 tat gtc ttg gct agt tac tcc ttg ttt act tgt tcc att gtt gaa gat  288
274 Tyr Val Leu Ala Ser Tyr Ser Leu Phe Thr Cys Ser Ile Val Glu Asp
275             85                  90                  95
277 gaa aaa aat aat ggg ggc caa aaa aga gtg tat ggt ttg tca caa gtg  336
278 Glu Lys Asn Asn Gly Gly Gln Lys Arg Val Tyr Gly Leu Ser Gln Val
279             100                 105                 110
281 gga aaa ttc ttt gtt aaa aat gaa aat ggt gca tca atg ggg cca ctt  384
282 Gly Lys Phe Phe Val Lys Asn Glu Asn Gly Ala Ser Met Gly Pro Leu
283         115                 120                 125
285 ttg gct ttg ctt caa aat aaa gta ttc ata aac agc tgg ttt gaa cta  432
286 Leu Ala Leu Leu Gln Asn Lys Val Phe Ile Asn Ser Trp Phe Glu Leu
```

VERIFICATION SUMMARY   DATE: 05/01/2002
PATENT APPLICATION: **US/09/937,204**   TIME: 11:41:49

Input Set : **A:\PTO.DC.txt**
Output Set: **N:\CRF3\05012002\I937204.raw**

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L:47 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:51 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:55 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:59 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:63 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:67 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:71 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:75 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:79 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:82 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:86 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:89 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:92 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:96 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:143 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:146 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:149 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:152 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:447 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:12
L:536 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:14
```